THE ANALYSIS OF VARIANCE (ANOVA)
for comparing multiple sample means (groups or treatments)
$\qquad$
$\mathrm{H}_{0}$ : The samples come from statistical populations with the same mean, i.e., $\mu_{\text {control }}=\mu_{\text {knee }}=\mu_{\text {eyes }}$.
$H_{A}$ : At least two samples come from different statistical populations with different means.

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## ANOVA

Research conclusion: Light treatment influences shifts in circadian rhythm.
How does light treatment influence shifts in circadian rhythm?

How do we know which group means differ from one another?

Why not simply not contrast
all pairs of means using a
two-sample mean t-test?
"The knees who say night"
Control vs. knee; control vs. eyes; knee vs. eyes?


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After ANOVA:

- Multiple testing and post hoc tests.
- The concept of family wise type I error and why we conduct ANOVAs first instead of twosample t-tests!


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Birthday and Preferences:
We were even able to observe an association between liking tea and liking eating vegetables (in a plausible direction) simply by separating individuals according to their birthdays.

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Another example of significance when there should be none Lee, K.L. et al. (1980) Clinical judgment and statistics. Lessons from a simulated randomized trial in
A simulated randomized clinical trial in coronary artery disease was conducted to illustrate the need for clinical judgment and modern statistical methods in assessing therapeutic claims in studies of complex diseases.

In this example, 1073 consecutive, medically treated coronary artery disease patients from the Duke University data bank were randomized into two groups. The groups were reasonably comparable and, as expected, there was no overall difference in survival between the two groups.
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Another example of significance when there should be none
1073 heart disease patients were RANDOMLY placed into two groups; no statistical difference was found in survival (not surprising given that they were randomly placed into groups as an exercise to demonstrate the issues with multiple testing) between the two groups.


11

Another example of significance when there should be none Lee, K.L. et al. (1980) Clinical judgment and statistics. Lessons from a simulated randomized trial in coronary artery disease. Circulation, 61: 508-515. Dol: 10.1166101.cir6.1.3.508
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But when patients were further subdivided into 18 prognostic categories, in a subgroup of 397 patients characterized by three-vessel disease and an abnormal left ventricular contraction, however, survival of group 1 patients was significantly different from that of group 2 patients. $\qquad$
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Another example of significance when there should be none
But when individuals between the two groups were then contrasted for their differences in survival according to 18 prognostic categories (heart morphology used to predict the likely outcome of a heart condition), for one of the categories the two groups differed in their survival. Any difference in survival should be due to chance alone as individuals were randomly divided into these categories.

Statistical tests across 18 prognostic categories


13

Another example of significance when there should be none
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But when patients were further subdivided into 18 prognostic categories, in a subgroup of 397 patients characterized by three-vessel disease and an abnormal left ventricular contraction, however, survival of group 1 patients was significantly different from that of group 2 patients.

Multivariable adjustment procedures revealed that the difference resulted from the combined effect of small imbalances in the distribution of several prognostic factors. Clinicians must exercise careful judgment in attributing such results to an efficacious therapy, as they may be due to chance or to inadequate baseline comparability of the groups.

14

Another example of significance when there should be none

- Patients grouped according as "three-vessel disease and an abnormal left ventricular contraction" were found to have differences between in survival between the two groups.
- However, patients were randomly assigned to each of the two groups in the beginning (i.e., survival versus non-survival).
How did that happen?

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Another example of significance when there should be none
Lee, K.L. et al. (1980) Clinical judgment and statistics. Lessons from a simulated randomized trial in
coronary artery disease. Circulation, 61:508-515. DOI: 10.1161/01.cir.61.3.508
1073 heart disease patients were randomly placed into two groups; no difference was found in survival (not surprising) between the two groups.
[akin to BIOL322 students divided according to their "birthdays"]
Individuals within each group were then contrasted according to 18 prognostic categories (heart morphology used to predict the likely outcome of a heart condition). [prognostics are akin to our 24 questions]

Individuals between the two groups were then contrasted for their differences in survival (any difference in survival should be due to chance alone as individuals were randomly divided into these categories). [p-values for a test comparing the two groups]

Patients grouped according as "three-vessel disease and an abnormal left ventricula contraction" were found to have differences between in survival between the two groups [students differ in their preferences for drinking tea and eating vegetables]

However, patients were randomly assigned to each of the two groups in the beginning (i.e., survival versus non-survival). [one should not expect differences related to odd/even birthdays]

How did that happen?
16


17


Why when comparing multiple means, one should start with an ANOVA and not by two-sample t-tests?


All the infinite $t$ values are possible under Ho , even the ones in the rejection region (they have a probability of alpha $=0.05$ to be sampled).
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If you conduct one or multiple tests, your probability of committing a type I error is still 0.05 (or another chosen alpha) $\qquad$
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From tutorial 9: There is a small statistical price to pay when inferring from samples: we need to accept a small risk of
$\qquad$ committing one sort of error [i.e., type I error] to avoid a bigger risk of committing another sort of error [i.e., type II error].


All the infinite $t$ values are possible under $\mathrm{H}_{0}$, even the ones in the rejection region (they have a probability of alpha $=0.05$ to be sampled).

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Why when comparing multiple means one should start with an ANOVA and not multiple $t$-test - because they inflate the number of false positive tests


If you conduct too many tests, you will eventually, by chance alone, hit a t value that is in the rejection region. Remember that the sampling distribution contains all infinite values for the $t$ statistic measuring the difference between two sample means assuming that $\mathrm{H}_{0}$ is true.

Because we eliminate implausible (Iow probability) values in the sampling distribution assuming the null hypothesis as true; assuming an alpha value to establish the rejection area, then it is obvious that if one conducts too many tests, one will eventually commit Type I errors for a given alpha (i.e., increase the number of false positive tests - reject when in reality one should not).

23

$\times$ *
If we set an alpha of 0.05 , i.e., acceptance area of $95 \%$ (0.95), then the chance of finding at least one significant test when you should not (i.e., false positive) out of 24 tests (groups) is: $\qquad$
$1-0.95^{24}=0.708$
$70.1 \%$ chance of finding at least 1 significant difference between odd and even born individuals in their preferences when $\mathrm{H}_{0}$ is true!

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Let's assume that 100 tests were conducted: $\qquad$

If we set an alpha of 0.05, i.e., acceptance area of 95\% (0.95), $\qquad$
then the chance of finding at least one significant test when you should not (i.e., false positive) out of 100 tests (groups) is:
$1-0.95^{100}=0.994$
$99.4 \%$ chance of finding at least 1 significant difference between group 1 and group 2 when $\mathrm{H}_{0}$ is true!

SO, $100 \%$ chance if you conduct 100 tests on samples that are expected to vary just due to chance alone (i.e., for which the null hypothesis $\mathrm{H}_{0}$ is true).

26

If we set an alpha of 0.05 , i.e., acceptance area of $95 \%$ (0.95), then the chance of finding at least one significant test when you should not (i.e., false positive) out of $\qquad$
1 test is obviously the original alpha:
$1-0.95^{1}=0.05$ $\qquad$
$5 \%$ chance of finding at least 1 significant test when $\mathrm{H}_{0}$ is true!


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If we set an alpha of 0.05, i.e., acceptance area of 95% (0.95), then the chance of
finding at least one significant lest when you should not (i.e., false positive) out of
24 tests is:
1-0.9544=0.708
70.1% chance of finding at least }1\mathrm{ significant test when }\mp@subsup{H}{0}{}\mathrm{ is true!
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The goal of performing an ANOVA before is to protect one against inflated type I errors due to $\qquad$ multiple pairwise testing.

When ANOVA is significant, which pairs of means can be "honestly" considered significant? $\qquad$
We need then a way to control for the possibility $\qquad$ of inflated type I errors due to multiple testing:
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The Tukey's honest test.

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$H_{A}$ : At least two samples come from different statistical populations with different means.

When ANOVA is significant, which pairs of means can be "honestly" considered significant?


31
(i.e., differences between means)?

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\binom{r}{2}=\frac{r!}{2!(r-2)!}=\frac{r(r-1)}{2}
$$


$\frac{3(3-1)}{2}=3$
\(\left.\begin{array}{l}Control-Knee <br>
Control-Eyes <br>

Knee-Eyes\end{array}\right]\) | 3 mean pairs |
| :---: |
| (contrasts) |

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32

The post-hoc (after ANOVA) - Tukey's honest test $\qquad$
There is a pair of hypotheses for each pair of means as follows: $\qquad$
$\mathrm{H}_{0}: \mu_{i}=\mu_{j}$ for each pair $i \neq j$
$\mathrm{H}_{\mathrm{A}}: \mu_{i} \neq \mu_{j}$ for each pair $\qquad$
$i$ and $j$ stand for the subscripts of the groups $\qquad$
(treatments) being compared.

| Control - Knee |  |
| :---: | :---: |
| Control-Eyes | O mean pairs |
| Knee - Eyes |  |

$\qquad$
$\begin{aligned} & \text { Control-Eyes } \\ & \text { Knee-Eyes }\end{aligned} \quad=\begin{gathered}3 \text { mean pairs } \\ \text { (contrasts) }\end{gathered}$ $\qquad$
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Tukey's honest test in R: we often use letters ( $a, b, c$., etc) to show on graphs the means that are different and similar. $\qquad$

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The test statistic for the Tukey Test is calculated as: $\qquad$
$\mathrm{Q}=\frac{\left|X_{i}-X_{j}\right|}{S E}$
$S E_{i-j}=\sqrt{\frac{s_{p(i, j)}^{2}}{2}\left(\frac{1}{n_{A}}+\frac{1}{n_{B}}\right)}$
$s_{p(i, j)}^{2}=\frac{d f_{i} s_{i}^{2}+d f_{j} s_{j}^{2}}{d f_{i}+d f_{j}}$
The quantity $s_{p}^{2}$ is called the pooled sample variance and is the average of the sample variances weighted by their degrees of freedom.

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ANOVA \& the Tukey-test:
Assumptions:

- Each of the samples (observations within groups) is a random sample from its population.
- The variable (shift in circadian rhythm) is normally distributed in each (treatment) population.
- The variances are equal among all statistical populations from which the treatments were sampled.

40

Testing differences in variances among populations - The Levene's test (too complex to understand its calculation for BIOL322 level but it is important to know its existence, utility and how to apply it in R). Hypotheses for the Levene's test:
$\mathbf{H}_{0}: \sigma_{\text {control }}^{2}=\sigma_{\text {knee }}^{2}=\sigma_{\text {eye }}^{2}$
$\mathbf{H}_{A}$ : At least one population variance ( $\sigma^{2}$ ) is different from another population variance or other population variances.

We need to generate evidence towards $\mathrm{H}_{0}$ to apply an ANOVA to the data at hands.

41

Testing differences in variances among populations - The Levene's test
$\mathbf{H}_{0}: \sigma_{\text {control }}^{2}=\sigma_{\text {knee }}^{2}=\sigma_{\text {eye }}^{2}$
$H_{A}$ : At least one population variance ( $\sigma^{2}$ ) is different from another population variance or other population variances.


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Levene's test:
Assumptions:

- Each of the samples (observations within groups) is a random sample from its population.
- The variable (shift in circadian rhythm) is normally $\qquad$ distributed in each (treatment) population.

43

Testing differences in variances among populations - The Levene's test (too complex for BIOL322 level but it is important to
$\qquad$ know its existence, utility and how to apply it in R). Hypotheses for the Levene's test

$\mathrm{P}=0.8545$. Based on an alpha $=0.05$, we should not reject the null hypothesis that: $\sigma_{\text {control }}^{2}=\sigma_{\text {knee }}^{2}=\sigma_{\text {eye }}^{2}$

Therefore, we should feel confident to conduct a standard ANOVA to the data (there is a Welch-like ANOVA).
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44

